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Search Results - Record(s) 1 through 2 of 2 returned.

	1.	<u>US20020007248A</u> .	Comparing	reference and	test DNA	samples	, useful in	forensic a	nalysis, by
deter	mir	ning a <u>likelihood rati</u>	o that allele	patterns matc	h. GILL, I	P, et al. C	C12Q001/6	8 G06F01	9/00.

2. <u>US20020009725A</u>. Determining defined source of DNA mixture by detecting allelic identity of DNA, determining probability functions for sample from different persons, using function ratio to determine loci ratios and combining them. <u>GILL</u>, P, et al. C12Q001/68.

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Term	Documents
LIKELIHOOD	99288
LIKELIHOODS	717
RATIO	1610144
RATIOS	269266
(1 AND (LIKELIHOOD ADJ RATIO)).USPT,EPAB,JPAB,DWPI.	2
(L1 AND LIKELIHOOD RATIO).USPT,EPAB,JPAB,DWPI.	2

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L3: Entry 1 of 2

File: DWPI

Feb 5, 2003

DERWENT-ACC-NO: 2002-041339

DERWENT-WEEK: 200310

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TITLE: Comparing reference and test DNA samples, useful in forensic analysis, by determining a likelihood ratio that allele patterns match

INVENTOR: GILL, P; BUCKLETON, J S; GILL, P D; WHITAKER, J P

PRIORITY-DATA: 2000GB-0009294 (April 15, 2000)

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PATENT-FAMILY:

PUB-NO	PUB-DATE		LANGUAGE	PAGES	MAIN-IPC
EP 1280933 A2	February 5,	2003	E	000	C12Q001/68
WO 200179541 A2	October 25,	2001	E	068	C12Q001/68
US 20020007248 A1	January 17,	2002		000	G06F019/00
AU 200148547 A	October 30,	2001		000	C12Q001/68

INT-CL (IPC): C12 Q 1/68; G06 F 19/00

ABSTRACTED-PUB-NO: US20020007248A

BASIC-ABSTRACT:

NOVELTY - A method of comparing reference DNA samples (A) with (part of) a test DNA (B), is new.

DETAILED DESCRIPTION - Alleles are identified for many loci in (B) to produce individual test results (ITR). These are compared with corresponding values (individual reference results, IRR) from an (A) and the probability (P) that IRR could lead by various routes to ITR, including routes where spurious information contributes to ITR, is determined. P for all loci are combined to give an expression for the probability that (B) matches (A), by calculating a likelihood ratio (LR).

INDEPENDENT CLAIMS are also included for the following:

- (a) method for calculating an LR for evaluating if (B) comes from a known source, or sources; and
- (b) method for investigating acceptable values for one or more variables related to DNA sample analysis.

USE - The method is used for forensic analysis.

ADVANTAGE - The method can be performed by non-experts; permits analysis of complex

samples; allows all information from analysis of DNA samples to be used in determination of results; allows quantification of potential spurious results; can validate approximations made in DNA analysis, and can be applied to results from small DNA samples.

ABSTRACTED-PUB-NO:

WO 200179541A EQUIVALENT-ABSTRACTS:

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CHOSEN-DRAWING: Dwg.0/10